

Exhibit A



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EMBOSS Align Results

Needle Results

Matrix: Blosum62
Open gap penalty: 10.0
Gap extension penalty: 0.5
Needle output: [needle-20070226-19371266036393.output](#)

[SUBMIT ANOTHER JOB](#)

```
#####  
# Program: needle  
# Rundate: Mon Feb 26 19:37:13 2007  
# Align_format: srspair  
# Report_file: /ebi/extserv/old-work/needle-20070226-19371266036393.output  
#####
```

```
#=====  
#  
# Aligned_sequences: 2  
# 1: EMBOSS_001  
# 2: EMBOSS_001  
# Matrix: EBLOSUM62  
# Gap_penalty: 10.0  
# Extend_penalty: 0.5  
#  
# Length: 524  
# Identity: 414/524 (79.0%)  
# Similarity: 451/524 (86.1%)  
# Gaps: 23/524 ( 4.4%)  
# Score: 2064.0  
#  
#  
#=====
```

018512-001420 us

Mouse SEQ 1

human SEQ 17

EMBOSS_001	1	MPMSSRDRDLHPGHHFGSCSPLSQLWPGP-EPKSVKGLYYSRARKVGNQ	49
EMBOSS_001	1	MPMPSRDGGHLPRHHYGHSPWSQLSSPMETPSIKGLYYRRVRKVGAL	50
EMBOSS_001	50	DASPEANLKEILVNVGGQRYLLPWSTLDAFPLSRLSRLRLCRSHEEITQL	99
EMBOSS_001	51	DASPVDLKKEILINVGGRRYLLPWSTLDRFPLSRLSKRLRLCRSYEEIVQL	100
EMBOSS_001	100	CDDYDEDSQEFFFDRNPSPAFGVIVSFLAAGKLVLLREMCALSFREELSYW	149
EMBOSS_001	101	CDDYDEDSQEFFFDRSPSPAFGVIVSFLAAGKLVLLQEMCALSFQEELAYW	150
EMBOSS_001	150	GIEETNLERCCLRKLLK---LEEAAELRREEA--AQRQQRQACHSEVQ	194
EMBOSS_001	151	GIEEAHLERCCLRKLLKLEEELEELAKLHREDVLRQQRETRRPASHS---	197
EMBOSS_001	195	ASRWASRMNQLREMVEDPQSGLPKGVFACLSVLFVATTAVSLCVSTMPDF	244
EMBOSS_001	198	-SRWGLCMNRLREMVENPQSGLPKGVFACLSILFVATTAVSLCVSTMPDL	246
EMBOSS_001	245	RAEEGKGECTRKCYIIFVVESICVAWFSLEFCLRFVQAPNKCQFFRGPLN	294
EMBOSS_001	247	RAEEDQGECSRKCYYIFIVETICVAWFSLEFCLRFVQAQDKCQFFQGPLN	296
EMBOSS_001	295	VIDILAISPYVSLAVSDESPEAGERPSSSSYLEKVGLVLRVLRALRILY	344

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